



**RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/016,869**

DATE: 04/16/2000  
TIME: 03:30:45

**INPUT SET: S35309.raw**

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46   (vii) PRIOR APPLICATION DATA:
47     (A) APPLICATION NUMBER: US 08/227,371
48     (B) FILING DATE: 14-APR-1994
49
50   (vii) PRIOR APPLICATION DATA:
51     (A) APPLICATION NUMBER: US 08/154,915
52     (B) FILING DATE: 18-NOV-1993
53
54   (vii) PRIOR APPLICATION DATA:
55     (A) APPLICATION NUMBER: US 07/991,997
56     (B) FILING DATE: 17-DEC-1992
57
58   (viii) ATTORNEY/AGENT INFORMATION:
59     (A) NAME: Vincent, Matthew P.
60     (B) REGISTRATION NUMBER: 36,709
61     (C) REFERENCE/DOCKET NUMBER: MIV-071.10
62
63   (ix) TELECOMMUNICATION INFORMATION:
64     (A) TELEPHONE: (617) 832-1299
65     (B) TELEFAX: (617) 832-7000
66
67

```

**ERRORED SEQUENCES FOLLOW:**

1071	(2) INFORMATION FOR SEQ ID NO:28:	
1072		
1073	(i) SEQUENCE CHARACTERISTICS:	
1074	(A) LENGTH: 434 base pairs	
1075	(B) TYPE: nucleic acid	
1076	(C) STRANDEDNESS: single	
1077	(D) TOPOLOGY: linear	
1078		
1079	(ii) MOLECULE TYPE: cDNA	
1080		
1081		
1082		
--> 1083	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:	
1084		
1085	CCCCATCGCG CCTTGGGANT GTGAGCNACC ATTGAGACTC ATNAATATAG CACTCGTTT	60   ord
1086		
1087	TCTTCTTGCA ACCCTGCCCN CCGCGCGGTC GCGCTNTCTC TGCCCTCCGC NGGGTGGACC	120   Sequence 2
1088		
1089	TGGAGCGAGC GCTTGAGCGG TCGGTCGGCG CNCCTGGANC AGCCAGGCGG GCAGTGGACT	180   missing.
1090		
1091	ACCTNCTGGA CCAGGGACCT GTGGGAGAGC GGTGNCGGCG GGTACATGCA CGTGAAGCCA	240
1092		
1093	TTGCGAGAAC TTTATCCATA AGTATTCAA TGCCGGTAGG GACGGCAAGA GAGGAGGGCG	300
1094		
1095	GGATGTNCCA CACATCTTG ACCTCAGGTT TCTAACGCCT GTTTCTTTC TGCCCTCTGC	360

*RECEIVED  
APR 24 2000  
TECH CENTER 1600/2900*

*Is this supposed to be 28?  
Also, Sequence 29 is  
missing. Most, parts of  
Sequence 28*

*(All next page)*

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1096		
1097	AGACATCCCC GATTGAAAGA ACCAGAGAGG CTCTGAGAAA CCTCCGGAAA CTTAGNTCAT	420
1098		
1099	CANTCGCCGN AAAAA	434
1100		
1101		

--> 1102 (2) INFORMATION FOR SEQ ID NO:30:

1103		
1104	(i) SEQUENCE CHARACTERISTICS:	
1105	(A) LENGTH: 502 base pairs	
1106	(B) TYPE: nucleic acid	
1107	(C) STRANDEDNESS: single	
1108	(D) TOPOLOGY: linear	
1109		
1110	(ii) MOLECULE TYPE: cDNA	
1111		
1112		
1113		

*This sequence directly follows above-mentioned sequence.*

1114	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:	
1115		

1116	AGAAATTAGA TCATCAGTCA CCGATCCTCC TACAGGGNCA CAACTGNCCC CGCCACAACC	60
1117		
1118	CACCCCGNTT TCGTAGTTTT CATTAGAAA ATAGAGCTTT TAAAAATGTC CTGCCTTTA	120
1119		
1120	ACGTAGATAT ATGCCTTCCC CCACTACCGN AAATGTCCAT TTATATCATN TTTTATATAT	180
1121		
1122	TCTTATAAAA ATGTAAAAAA GAAAAACACC GCTTCTGCCT TTTCACTGTG TTGGAGTTTT	240
1123		
1124	CTGGAGTGAG CACTCACGCC CTAAGCGCAC ATTCACTGTGG GCATTCTTG CGAGCCTCGC	300
1125		
1126	AGNCTCCGGA AGCTGTCGAC CTCGAGGGGG GGNCCGGTAC CCAATTGCC CTATAGTGAG	360
1127		
1128	TCGTATTACA ATTCACTGGN CGNCNTTTT ACAACGTCGG TGGACTGGGA AAACCCGGN	420
1129		
1130	GTTACCCAAC TTTAATCGNC TTGGAGGACA TCCCCCTTTT CGCCAGNTGG GGTTATAGNG	480
1131		
1132	AAGAGGGCCN CACCNNTCGC CC	502
1133		
1134		

1236 (2) INFORMATION FOR SEQ ID NO:34:

1237		
1238	(i) SEQUENCE CHARACTERISTICS:	
1239	(A) LENGTH: 88 amino acids	
1240	(B) TYPE: amino acid	
1241	(C) STRANDEDNESS:	
1242	(D) TOPOLOGY: linear	
1243		
1244	(ii) MOLECULE TYPE: peptide	
1245		
1246		
1247	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:	

*OK*

**RAW SEQUENCE LISTING  
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1248  
1249  
1250  
1251 Ile Gln Val Met Met Gly Ser Ala Arg Val Ala Glu Leu Leu Leu Leu  
1252 1 5 10 15  
1253 His Gly Ala Glu Pro Asn Cys Ala Asp Pro Ala Thr Leu Thr Arg Pro Val  
1254 20 25 30  
1255  
1256 His Asp Ala Ala Arg Glu Gly Phe Leu Asp Thr Leu Val Val Leu His Arg  
1257 35 40 45 50  
1258  
1259 Ala Gly Ala Arg Leu Asp Val Arg Asp Ala Trp Gly Arg Leu Pro Val Asp  
1260 55 60 65  
1261  
1262 Leu Ala Glu Glu Arg Gly His Arg Asp Val Ala Gly Tyr Leu Arg Thr Ala  
1263 70 75 80 85  
1264  
1265  
1266 Ala Gly Asp

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PAGE: 1

**SEQUENCE VERIFICATION REPORT  
PATENT APPLICATION US/09/016,869**

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***INPUT SET: S35309.raw***

Line	Error	Original Text
13	Number of Sequences (34) Doesn't Equal Actual Count (33)	(iii) NUMBER OF SEQUENCES: 34
1083	Wrong Sequence Number	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:
1102	Sequence 29 missing	(2) INFORMATION FOR SEQ ID NO:30: